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2018-12

Altinci , P , Mutluay , M , Tjäderhane , L & Tezvergil-Mutluay , A 2018 , ' Effect of calcium fluoride on the activity of dentin matrix-bound enzymes ' , Archives of Oral Biology , vol. 96 , pp. 162-168 . <https://doi.org/10.1016/j.archoralbio.2018.09.006>

<http://hdl.handle.net/10138/307704>

<https://doi.org/10.1016/j.archoralbio.2018.09.006>

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Effect of calcium fluoride on the activity of dentin matrix-bound enzymes

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ARTICLE INFO

Keywords:

Dentin matrix
CaF₂
Collagen degradation
Cysteine cathepsins
MMPs

ABSTRACT

Objective: Matrix metalloproteinases (MMPs) and cysteine cathepsins (CCs) are two distinct enzymatic pathways responsible for the degradation of collagen fibrils in demineralized dentin. NaF and KF have been shown to inhibit salivary MMP-2, -9 and CCs. This study investigated the inhibitory effect of calcium fluoride (CaF₂) on the dentin matrix-bound MMPs and CCs.

Design: Phosphoric acid (10%)-demineralized dentin beams (1 × 2 × 6 mm) were incubated at 37 °C in an 1 ml of artificial saliva (AS, control), or AS with 6, 12, 24, 48, 120, 179 and 238 mM F containing CaF₂ (n = 10/group) for 1, 7 and 21 days. All groups were further incubated in AS only for 6 months. Total MMP activity, dry mass loss, CTX and hydroxyproline (HYP) analyses were performed after each incubation. The beams were examined under scanning electron microscopy (SEM). MMP-2 and MMP-9 activities were screened with gelatin zymography. Data were analyzed by using ANOVA and Tukey HSD tests (p = .05).

Results: The total MMP activity was similar for all groups after 21 days and 6 months. After 21 days, the cumulative mass loss and CTX levels were lower compared to control for the CaF₂ ≥ 48 and CaF₂ ≥ 120 mM, respectively (p < .05). After 6 months, no significant difference was detected in the dry mass loss and CTX compared to the control (p > .05), whereas HYP level was higher with F 24 and 238 mM groups. CaF₂-like minerals were observed on the beams under SEM. There was no gelatinase inhibition in zymography.

Conclusion: CaF₂ does not prevent the degradation of demineralized dentin matrices due to the catalytic activity of MMPs and CCs.

1. Introduction

Auto-degradation of demineralized dentin organic matrix occurs by the catalytic activity of the endogenous dentin enzymes including mainly matrix metalloproteinases (MMP)-2, -8, -9 and cysteine cathepsin (CC)-K (Chaussain et al., 2013; Chaussain-Miller, Fioretti, Goldberg, & Menashi, 2006; Frassetto et al., 2016; Mazzoni et al., 2015; Tjäderhane et al., 1998; Tjäderhane, 2015; Tjäderhane, Buzalaf, Carrilho, & Chaussain, 2015). It is thought that these enzymes physiologically function during the teeth development phase, and fossilized in dentin with the maturation of teeth (Chaussain-Miller et al., 2006). Many studies showed that, due to the demineralization of dentin during cariogenic and erosive processes as well as adhesive bonding procedures requiring the use of acid-etchant or acidic resin monomers, these enzymes become re-activated, and cut off the type I collagen fibrils constituting dentin organic matrix (Buzalaf, Charone, & Tjäderhane,

2015; Chaussain-Miller et al., 2006; Mazzoni et al., 2015; Tjäderhane et al., 2013a). As a consequence, remineralization cannot take place in these demineralized zones, leading to progressive dentin tissue loss (Tjäderhane et al., 1998). At the adhesive interfaces, these enzymatic activities together with the hydrolysis of resin components cause the premature failure of bonded restorations due to microleakage and secondary caries (Buzalaf et al., 2015; Tjäderhane et al., 2013a; Tjäderhane, 2015). Inhibition of dentin matrix-bound MMPs and CCs may hinder the progression of early carious and erosive lesions, and can enhance the clinical longevity of adhesive restorations by improving the hydrolytic stability of adhesive interfaces (Mazzoni et al., 2015; Tjäderhane et al., 2013a, 2013b; Tjäderhane, 2015; Tjäderhane et al., 2015).

Fluoride (F) is an antimicrobial anion improving the tooth demineralization resistance by forming less soluble fluorhydroxyapatite and fluorapatite minerals by interacting with the Ca²⁺ of enamel and dentin

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<https://doi.org/10.1016/j.archoralbio.2018.09.006>

Received 28 February 2018; Received in revised form 6 August 2018; Accepted 9 September 2018

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(ten Cate, 2013). Fluoridated anti-demineralization agents create CaF_2 or CaF_2 -like crystals on teeth surfaces, so-called F reservoirs, releasing F ions to the environment under acidic challenges (Buzalaf, Pessan, Honório, & ten Cate, 2011; Lussi, Hellwig, & Klimek, 2012; Ogaard, 2001). During demineralization, the released F ions intervene to the cariogenic process, and promote remineralization. Owing to these benefits, CaF_2 has been experimentally incorporated into the composite resins in order to formulate F-releasing and antimicrobially effective restorative materials to prevent secondary caries (Xu, Moreau, Sun, & Chow, 2010, 2010b; Cheng et al., 2012; Weir et al., 2012).

Recent studies have shown that F ions can inhibit the catalytic activity of salivary MMPs (Kato et al., 2014) as well as dentin matrix-bound CCs (Altinci et al., 2016). Moreover, NaF and KF demonstrated quite similar inhibitory effect on CCs (Altinci et al., 2016; Altinci, Mutluay, Tjäderhane, & Tezvergil-Mutluay, 2017). Hence, this study aimed to investigate if CaF_2 has anti-enzymatic effect on the dentin matrix-bound MMPs and CCs, preventing the degradation of demineralized dentin matrices. The tested null hypothesis was CaF_2 does not prevent the proteolytic degradation of demineralized dentin matrices due to the dentin matrix-bound MMPs and CCs.

2. Materials and methods

Fifty sound human molars stored in 0.02% NaN_3 disinfectant solution were used for the study series according to the approval of Ethical Committee, Faculty of Medicine, University of Oulu (#19/2006). Two horizontal sections were done from the mid-coronal portions of the teeth to obtain dentin disks using a low-speed diamond saw (Isomet saw, Buehler Ltd., Lake Bluff, IL, USA) under continuous water-cooling. The disks were further sectioned into dentin beams (1 mm × 2 mm × 6 mm). The beams were demineralized in 10% phosphoric acid for 24 h, rinsed in distilled water under constant stirring at 4 °C for 8 h, and dried to a constant weight in a vacuum desiccator containing dry silica beads. After measuring the dry masses with a digital microbalance with a sensitivity of 0.001 mg (Mettler XP6 Microbalance, Mettler Toledo, Hightstown, NJ, USA), each beam was allocated to a group based on similar baseline dry mass. Then, the beams were rehydrated in distilled water at 4 °C for 2 h, placed in labeled polypropylene tubes for incubation (n = 10/group).

The incubation was performed in 1 ml of artificial saliva (AS, 5 mM HEPES, 2.5 mM $\text{CaCl}_2 \cdot \text{H}_2\text{O}$, 0.02 mM ZnCl_2) with CaF_2 in the F concentrations of 6, 12, 24, 48, 120, 179, 238 mM at pH 7.4 for 1, 7 or 21 days. The control group beams were incubated only in AS. Afterwards, all groups were incubated only in AS without any F content at pH 7.4 for 6 months. Due to the low water solubility of CaF_2 , the incubation solution was prepared by heating up to 60 °C in a beaker under constant stirring, and then cooled to room temperature. The F content of the solution was directly measured using a fluoride ion-selective electrode (DX219-F, Mettler Toledo, Urdorf, Switzerland). A shaking bath (OLS200, Grant Instruments, Cambridge, UK) was used for the incubation at 60 cycles/min and at 37 °C under constant pH. At the end of each incubation period, the medium was stored at –70 °C until further analysis of collagen degradation products.

2.1. Total MMP activity

A generic colorimetric MMP assay (Sensolyte Generic MMP assay; Anaspec, San Jose, CA, USA) was used to measure the total activity of dentin matrix-bound MMPs subjected to CaF_2 (Altinci et al., 2016; Ozcan et al., 2015). Demineralized dentin beams (phosphoric acid 10%) used as enzyme source were immersed in 300 µl of chromogenic thiopeptide substrate and assay buffer for 60 min at 25 °C (n = 10/group). Then, the beams were removed from the wells, and the absorbance values were measured at 412 nm. Based on the similar baseline MMP activity, the beams were divided into groups (n = 10/group). At the end of each incubation period, the beams were rinsed with distilled

water, and their total MMP activities were measured again. The results were calculated as a percentage of the untreated baseline level to determine the percent of inhibition or activation.

2.2. Dry mass loss

As an indirect indicator of dentin matrix degradation, the dry mass loss of the beams was measured at the end of each incubation period (n = 10/group). For this purpose, the buffer salts were washed off in distilled water under constant stirring at 4 °C for 8 h. Then, the beams were dried to a constant weight in the desiccator for 72 h. After weighing the beam dry masses with the digital microbalance, they were rehydrated in distilled water at 4 °C for 2 h, and taken to the next incubation period with a fresh incubation medium. The results were calculated based on the respective baseline dry masses, and expressed as the rate of dry mass loss.

2.3. Hydroxyproline analysis

Approximately 9.6 mass% of type I collagen fibrils is constituted by an amino acid hydroxyproline (HYP). The measurement of HYP quantity in the incubation medium of dentin beams indicates the total liberated collagen by the activity of both MMPs and CCs (Tezvergil-Mutluay, Agee, Hoshika, Tay, & Pashley, 2010). To hydrolyze the collagen fragments into amino acids, 400 µl of incubation medium of dentin beams was mixed with 400 µl HCl to a final acid concentration of 6 N HCl in glass ampules (Wheaton, Millville, NJ, USA) (n = 10/group). The ampules were sealed using a flame sealer (Ampulmatic, Bioscience, Inc., Allentown, PA, USA), and incubated at 118 °C for 18 h. After cooling to room temperature, the content was dried in the desiccator connected to a NaOH pellet container to trap HCl vapor, and processed according to the hydroxyproline method described by Jamall, Finelli, and Que Hee, (1981). To each ampule, 50% isopropanol in water (1.2 ml), and Chloramine T solution with acetate citrate buffer (0.2 ml) were pipetted. After 10 min., Ehrlich's solution (1 ml) prepared with 4-dimethylaminebenzaldehyde, 60% perchloric acid and 100% isopropanol was added. Following incubation at 50 °C for 90 min., the liquids were transferred to the cuvettes, and the absorbances were read by a spectrophotometer (Model UV-A180, Shimadzu, Tokyo Japan) at 558 nm. The results were calculated as µg of hydroxyproline/mg of the dry mass of demineralized dentin before incubation.

2.4. CTX analysis

C-terminal crosslinked telopeptide of type I collagen (CTX) is produced by the catalytic activity of CCs. The CTX levels in the incubation medium of dentin beams can be regarded as an indicator of the CC activity in demineralized dentin matrix (Garnero et al., 2003; Tezvergil-Mutluay et al., 2015). Therefore, the CTX levels of the aliquots (20 µl) were measured with specific assay kits for human serum CTX (Serum CrossLaps ELISA, Immunodiagnostic System, Farmington, UK) using a plate reader (Synergy HT, BioTek Instruments, Vermont, USA) (n = 10/group). The absorbances were measured at 450 nm with 650 nm as reference. The experiments were performed in triplicate.

2.5. Scanning electron microscopy and EDS analysis

From each group, 5 representative beams were gold-sputtered (SC7620 Sputter Coater, Quorum, East Sussex, UK) after 21 days and 6 months incubation, and scanning electron microscopy (SEM) images were taken with a SEM device (Phenom-World, Eindhoven, The Netherlands) operating at 10 kV. The beams were fractured after kept in liquid nitrogen for 5 min, and further examined for elemental composition with an EDS detector integrated to SEM device at 15 kV with line scan function up to approximately 100 µm in depth of the beams.

2.6. Gelatin zymography

Gelatin zymography for the MMP-9 and MMP-2 activities in dentin was performed using dentin powder as enzyme source. For this purpose, coronal dentin pieces were obtained from approximately 100 sound molar after removing the roots, coronal enamel and intrapulpal dentin with high-speed burs under water cooling. Dentin pieces were placed into the metal containers of a cryomilling machine together with 2 metal balls. After the metal lids were tightened, the container was dipped into the liquid nitrogen for 5 min, then placed to the machine, and shaken for 2.5 min. The resultant fine mineralized dentin powder was collected in tubes, and stored at -70°C until use.

Gelatin zymography was performed according to the protocol of Mazzoni et al. (2013). Demineralized (10% phosphoric acid for 10 min) dentin powder batches (200 mg/group) were treated with CaF_2 in the F concentrations of 6, 12, 24, 48, 120, 179, 238 mM (100 μl) for 5 min at 4°C , centrifuged (12,000 g, 20 min), and the supernatants were discarded. Following washing with distilled water, the samples were suspended in extraction buffer containing 50 mM Tris-HCl pH 6, containing 5 mM CaCl_2 , 100 mM NaCl, 0.1% Triton X-100, 0.1% nonionic detergent P-40, 0.1 mM ZnCl_2 , 0.02% NaN_3 for 24 h at 4°C , sonicated (for 10 min, twice), and centrifuged (12,000 g, 20 min at 4°C). The supernatants were collected, and concentrated (10,000 g, 20 min at 4°C) to 50 μl total volume via Vivaspin centrifugal concentrator (10,000 kDa cut off; Vivaspin Sartorius Stedim Biotech, Goettingen, Germany). Total protein concentration of the dentin extracts was determined by Bradford assay. Then, the protein aliquots of 100 μg were diluted (1/4 ratio) in Laemmli sample buffer. After the electrophoresis performed in 10% sodium dodecyl sulfate-polyacrylamide gel (SDS-PAGE containing 1 mg/ml fluorescein-labeled gelatin), and washing in 2% Triton X-100 for 1 h, the gel was incubated in zymography activation buffer (50 mmol/L Tris-HCl, 5 mmol/L CaCl_2 , pH 7.4) for 48 h. Prestained low-range molecular weight SDS-PAGE standards (Bio-Rad, Hercules, CA, USA) were used as molecular-weight markers. The gelatinase activity in the demineralized dentin powder was also analyzed. The results were monitored using a long-wave UV light scanner (Chemidoc Universal Hood, Bio-Rad, Hercules, CA, USA). The entire experiment was performed in duplicate.

2.7. Statistical analysis

Data of total MMP activity, dry mass loss, HYP and CTX analyses were evaluated with Shapiro-Wilk and Brown-Forsythe test for normality and equal-variance assumptions, respectively. Since these parameters were satisfied, repeated measures ANOVA tests were applied based on 'pretreatment' as group variable, and 'timepoint' as the repeated factor. In case significant interactions were found, one-way ANOVA with Tukey HSD *post hoc* tests were performed. All statistical tests were performed using SPSS Statistics 23.0 software (IBM Corporation, Armonk, NY, USA) at $p = .05$.

3. Results

The mean (\pm SD) total MMP activity of dentin beams after each incubation period is shown in Fig. 1. After 1 day of incubation, there was a reduction in the MMP activity of CaF_2 groups, which was only significant for F 238 mM group ($p < .05$). After 7 days of incubation, there was a gradual decrease in the MMP activities of F groups, which were statistically significant for F 24, 179 and 238 mM. However, after 21 days of incubation, there were no significant differences between the MMP activities of F groups and the control ($p > .05$).

The mean (\pm SD) dry mass loss of the control and the CaF_2 groups after 1, 7 and 21 days of incubation is presented in Fig. 2. Compared to the control, the dry mass loss of the F groups significantly decreased starting from the 24 mM F group after 1 day, and 48 mM F after the 7 days of incubation. As seen in Fig. 3, in consistence with dry mass loss

results, there were slight decreases in the CTX levels, which was statistically significant for $F \geq 120$ mM groups after 21 days. However, no significant difference was detected between the HYP levels of the control and F groups at all timepoints ($p > .05$).

The total MMP activity, dry mass loss, hydroxyproline and CTX assay results of the control and CaF_2 groups incubated for 6 months in AS are shown in Fig. 4. The MMP activity ranged between 45–60% among the groups with no significant difference ($p > .05$). Likewise, there were no significant differences between the dry mass loss and the CTX levels of CaF_2 groups and the control ($p > .05$). The HYP levels of F 24 and 238 mM were significantly higher than the control ($p < .05$).

The SEM-EDS images of the representative beams after the 21 days and 6 months of incubation are shown in Fig. 5. As confirmed by EDS, CaF_2 -like minerals were observed on dentin beams with increasing density in correlation with the increasing CaF_2 concentrations. After 6 months incubation, the precipitations were still dense over the beams. Inside the dentin, F ion infiltration was confirmed to the approximately 100 μm in depth.

Fig. 6 shows the gelatin zymography results of demineralized dentin powder samples treated with increasing concentrations of CaF_2 with F 6–238 mM. In the protein extract of demineralized dentin powder the bands detected at 95 kDa and 68–72 kDa, corresponding to the molecular weights of proMMP-9, and active and proforms of MMP-2, respectively. The bands showing MMP-2 and -9 activities in CaF_2 groups were remarkably noticeable, indicating that there was no inhibition.

4. Discussion

This study investigated the effect of CaF_2 on the catalytic activity of dentin matrix-bound MMPs and CCs for the prevention of demineralized dentin matrix degradation. As direct indicators of the MMP activity, total MMP activity assay and gelatin zymography were performed. Despite the gradual decrease in the total MMP activity after 7 and 21 days of incubation, there was no inhibition in the MMP-2 and MMP-9 activities according to the zymography. Hence, the indirect indicators of the enzymatic degradation in dentin organic matrix were analyzed involving the beam mass loss, and CTX and HYP assays. Up to 21 days of incubation, mass loss and CTX results consistently indicated that CaF_2 could be effective in the inhibition of CCs. However, there was no decrease in the HYP levels of CaF_2 groups compared to control at all incubation periods. Furthermore, there was no CC inhibition according to the CTX results after 6 months of incubation. Therefore, it was concluded that CaF_2 does not prevent the degradation of demineralized dentin organic matrices due to MMPs or CCs. Hence, the tested null hypothesis was accepted.

It has been previously suggested that salivary MMPs and dentin matrix-bound CCs can be inhibited by the high levels of NaF and KF (Altinci et al., 2016; Kato et al., 2014; Mei et al., 2014). Moreover, KF was effective in the long-term inhibition of CCs (Altinci et al., 2017). These results have been attributed to highly electronegative F ions, binding to the zinc and calcium in the enzyme structure, thereby blocking the catalytic domain, or causing conformational changes in the enzyme morphology. In the present study, CaF_2 was used as fluoride source, however similar inhibitory effects could not be obtained. One reason might be the low water solubility of CaF_2 as a difluoride compound compared to the highly water-soluble monofluorides, NaF and KF. Another reason might be the substrate preference of F ions. Assuming that F ions primarily prefer binding to Ca^{2+} ions, the interaction of F compounds with dentin matrix-bound MMPs and CCs can be explained. In case of no free Ca^{2+} ions in the media, F ions can bind to structural Ca^{2+} and Zn^{2+} ions of MMPs, which clarifies the inhibition of salivary MMPs with high F concentrations (Kato et al., 2014). On the other hand, there were Ca^{2+} ions in artificial saliva used in this study, readily available for F binding. After reacting with these Ca^{2+} ions, the remaining F ions can bind to the enzymes. At this point, it can be also suggested that the required F concentration could be lower for

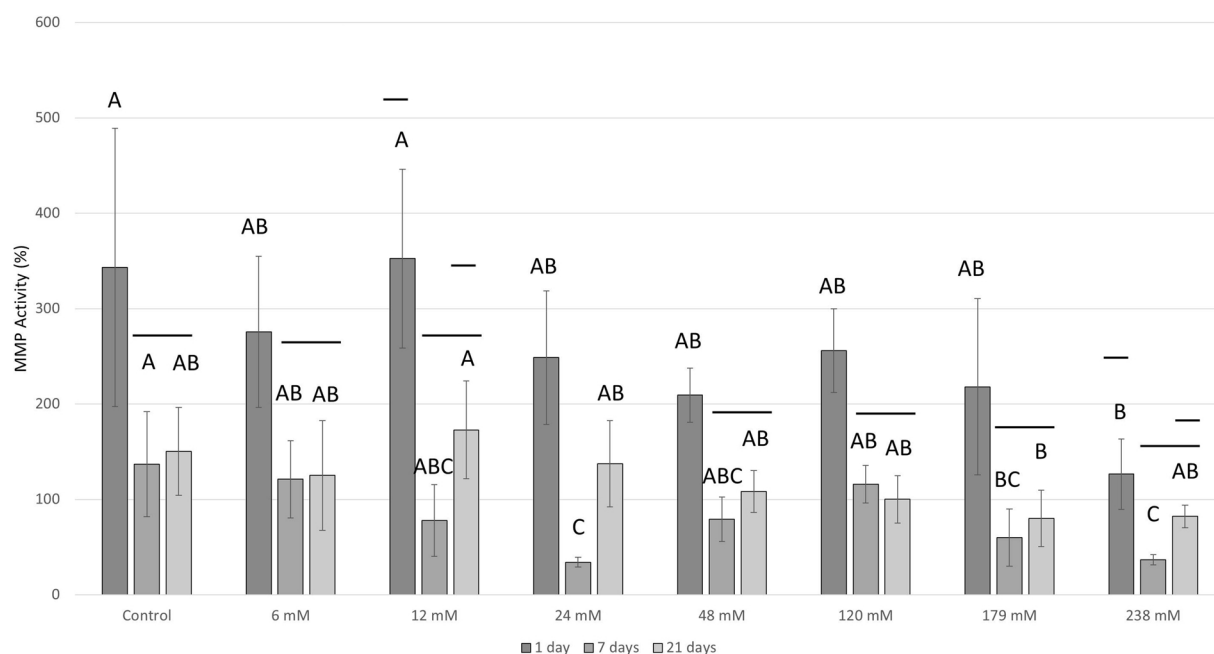


Fig. 1. Total MMP-activity rates in the demineralized dentin beams after incubated for 1, 7, 21 days in the incubation medium with the increasing F concentrations (6–238 mM) or without F (control). The MMP-activity rate of each beam was cumulatively calculated as a percentage of the baseline MMP-activity measured before incubations. For each time period, the groups with the same uppercase letters are not statistically significant ($p > .05$). For each fluoride concentration as well as the control, time period columns connected by a solid black bar are not statistically significant ($p > .05$).

cathepsin inhibition than MMPs since dentin matrix-bound CCs were significantly inhibited by NaF and KF (Altinci et al., 2016, 2017). Higher F concentrations also resulted in denser mineral precipitations on dentin, possibly because the initial precipitations behaved as nuclei for crystal growth (Lussi et al., 2012). Therefore, MMP inhibition could not be observed in demineralized dentin with high F concentrations either.

CaF₂-like minerals can form on dentin depending on the solubilization of Ca²⁺ ions from tooth surface, F exposure duration, F concentration and the pH of the F solution (Ogaard, 2001). Present study was implemented at pH 7.4 considered the optimum pH range for the activity of MMPs and the structural stability of collagen fibrils. On the other hand, CCs demonstrate higher catalytic activity at pH 5.5 (Tezvergil-Mutluay et al., 2015). If the same study was performed at pH

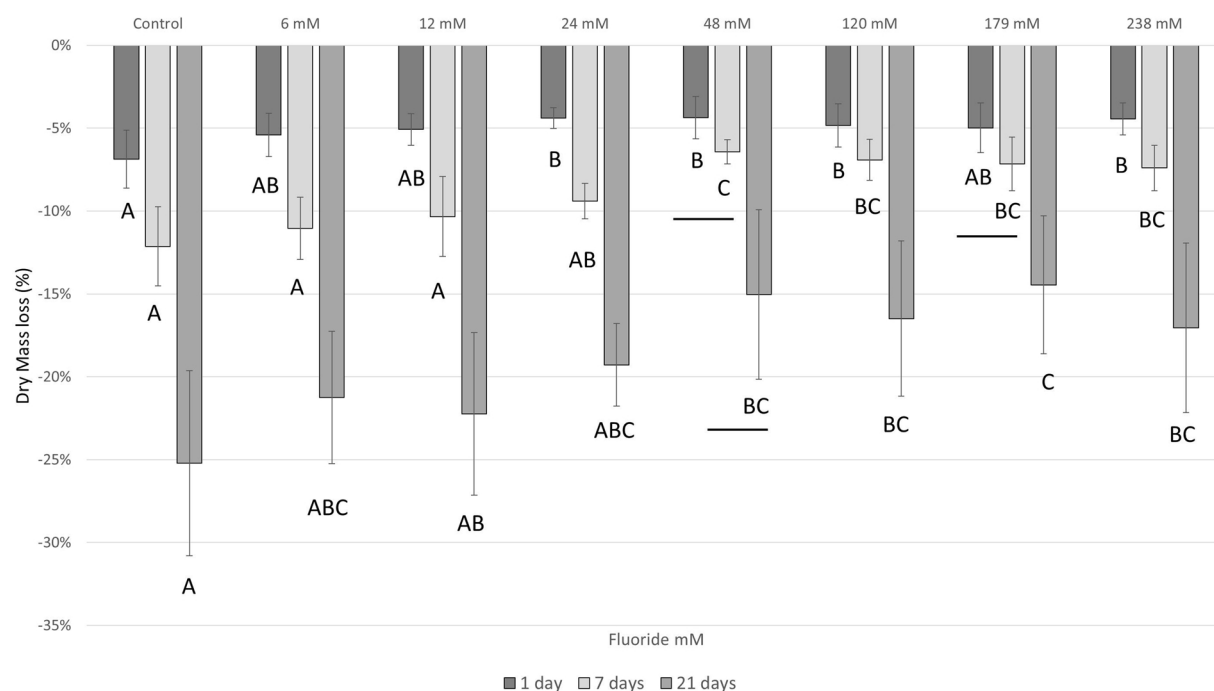


Fig. 2. Loss of cumulative dry mass of demineralized dentin beams after incubated for 1, 7, 21 days in the incubation medium with the increasing F concentrations (6–238 mM) or without F (control). The loss of dry mass from each beam was calculated as a percentage of the original dry mass of that beam at baseline. For each time period, the groups with the same uppercase letters are not statistically significant ($p > .05$). For each fluoride concentration as well as the control, time period columns connected by a solid black bar are not statistically significant ($p > .05$).

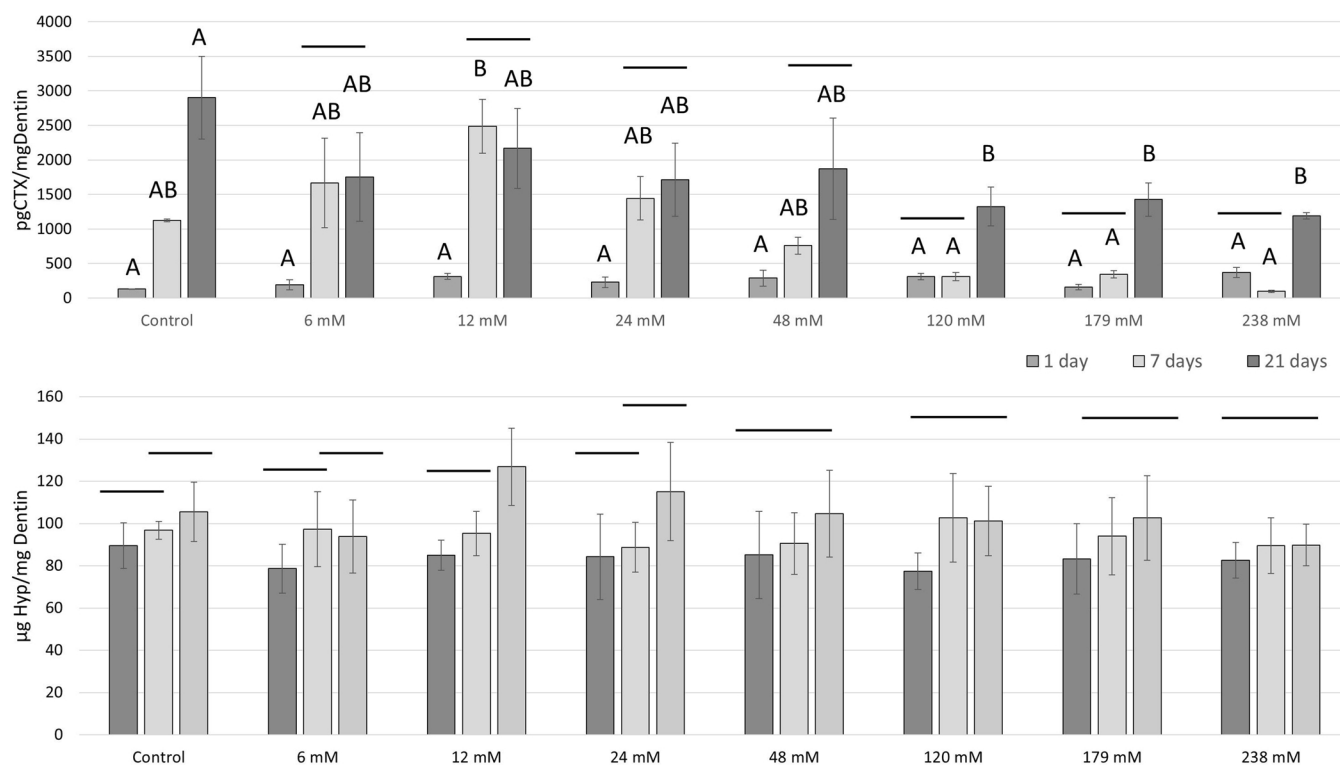


Fig. 3. Bar charts comparing the liberated mean CTX and hydroxyproline levels with standard deviations from demineralized dentin beams after incubated for 1, 7, 21 days in the incubation medium with the increasing F concentrations (6–238 mM) or without F (control). For each time period, the groups with the same uppercase letters are not statistically significant ($p > .05$). For each fluoride concentration as well as the control, time period columns connected by a solid black bar are not statistically significant ($p > .05$).

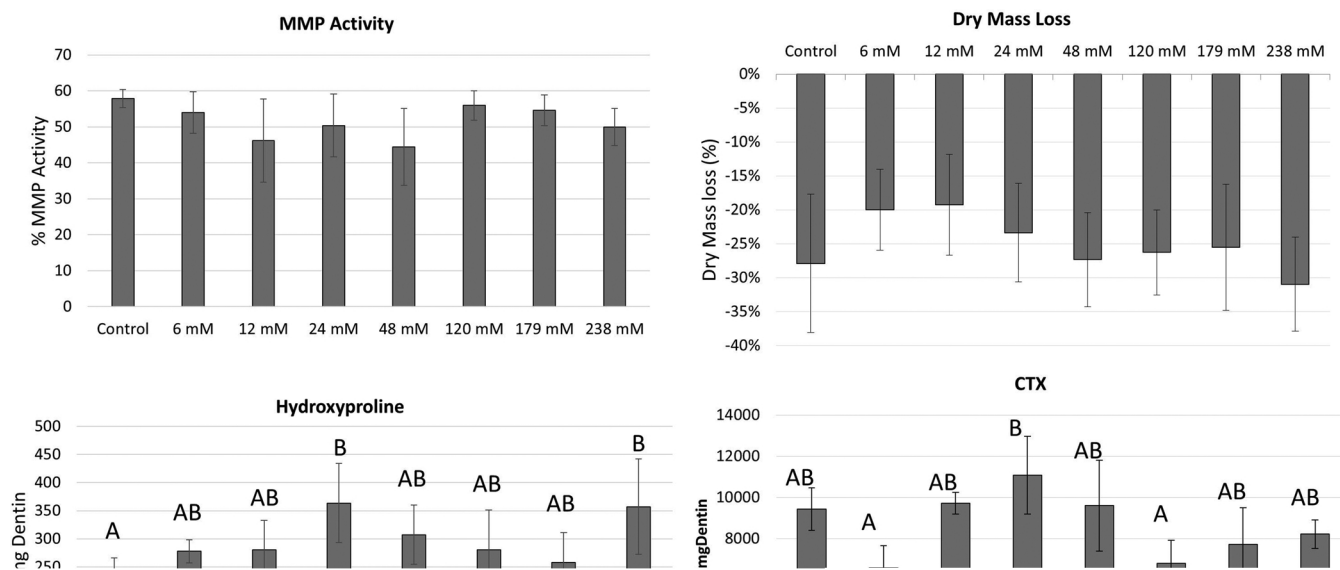


Fig. 4. Bar charts of total MMP activity, dry mass loss, hydroxyproline and CTX assay results obtained after the demineralized dentin beams of control and F (6–238 mM) groups were incubated in AS for 6 months. The statistical significance between the hydroxyproline level of 24 mM F and control groups was marked with an asterisk (*) ($p < .05$).

ions with the enzymes.

Author contributions

P. Altinci, M. Mutluay, L. Tjäderhane and A. Tezvergil-Mutluay contributed to the research design. P. Altinci was responsible from the laboratory testing and data gathering. P. Altinci, M. Mutluay, L. Tjäderhane and A. Tezvergil-Mutluay analyzed the data. P. Altinci drafted the manuscript. All authors participated in the editing and final preparation of the manuscript.

Acknowledgements

This study was funded by the grant #296653 from the Academy of Finland to AT-M (PI) and EVO funding #13140 of Turku University Hospital to AT-M (PI). The funders had no role in study design, data collection and analysis, decision to publish, or preparation of the manuscript. The authors do not have a financial interest in products, equipment, and companies cited in the manuscript. The authors thank the staff of the Helsinki Finnish Student Health Service (FSHS) and Turku Student Health Service for collecting the teeth for the experiments.

The authors would like to thank to the biomedical laboratory technician Katja Sampalahti for performing CTX assays, to Docent Dr. Roda Seseogullari-Dirihan for her contributions to the total MMP activity assays, and to Kaveh Nik Jamal, MSc for performing SEM-EDS examinations in this study.

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